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SEQUENCE LISTING

<110> [redacted] Atsuhiko

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Nihira, Takuya

Shindo, Takuya

<120> METHOD FOR INDUCTION OF GENE EXPRESSION IN PLANT AND PLANT TREATED THEREBY

<130> 5405/18

<140> 10/049,710

<141> 2002-02-15

<150> PCT/JP01/05096

<151> 2001-06-15

<150> JP 2000-180466

<151> 2000-06-15

<160> 11

<170> PatentIn version 3.1

<210> 1

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<212> DNA

<213> Streptomyces virginiae

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<221> CDS

<222> (1)..(699)

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<301> Okamoto, S., Nakamura, K., Nihira, T. and Yamada, Y.

<302> Virginiae butanolide binding protein from Streptomyces virginiae. Evidence that VbrA is not the virginiae butanolide binding protein and re-identification of the true binding protein

<303> Journal of Biological Chemistry

<304> 270

<305> 20

<306> 12319-12326

<307> 1995-05-19

<308> D32251

<309> 1994-07-19

<313> (1)..(699)

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<301> Okamoto, S., Nakamura, K., Nihira, T. and Yamada, Y.

<302> Virginiae butanolide binding protein from Streptomyces virginiae. Evidence that VbrA is not the virginiae butanolide binding protein and re-identification of the true binding protein

<303> Journal of Biological Chemistry

<304> 270

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<307> 1995-05-19

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<309> 1994-07-19

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cgc acg cgg cag gcg atc gtg cgg gca gcc gcc tcg gtc ttc gac gag       96
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Arg Thr Arg Gln Ala Ile Val Arg Ala Ala Ala Ser Val Phe Asp Glu
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tac ggg ttc gag gcc gcc aca gtg gca gag atc ctc tcg cgg gcc tcg      144
Tyr Gly Phe Glu Ala Ala Thr Val Ala Glu Ile Leu Ser Arg Ala Ser
        35                  40                  45

gtc acc aag ggc gcg atg tac ttc cac ttc gct tcc aag gaa gag ctg      192
Val Thr Lys Gly Ala Met Tyr Phe His Phe Ala Ser Lys Glu Glu Leu
    50                  55                  60

gcc cgc ggc gtg ctg gcc gag cag acc ctg cac gtg gcg gtg ccg gaa      240
Ala Arg Gly Val Leu Ala Glu Gln Thr Leu His Val Ala Val Pro Glu
65                  70                  75                  80

tcc ggc tcc aag gcg cag gaa ctg gta gac ctc acc atg ctg gtc gcc      288
Ser Gly Ser Lys Ala Gln Glu Leu Val Asp Leu Thr Met Leu Val Ala
                85                  90                  95

cac ggc atg ctg cac gat ccg atc ctg cgg gcg ggc acg cgg ctc gca      336
His Gly Met Leu His Asp Pro Ile Leu Arg Ala Gly Thr Arg Leu Ala
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ctg gac cag ggg gcg gtg gac ttc tcc gac gcc aac ccg ttc ggc gag      384
Leu Asp Gln Gly Ala Val Asp Phe Ser Asp Ala Asn Pro Phe Gly Glu
        115                 120                 125

tgg ggc gac atc tgc gcc cag ctc ctg gcg gag gca cag gaa cgg ggg      432
Trp Gly Asp Ile Cys Ala Gln Leu Leu Ala Glu Ala Gln Glu Arg Gly
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gag gtg ctt ccg cac gtg aac ccg aaa aag acc ggc gac ttc atc gtc      480
Glu Val Leu Pro His Val Asn Pro Lys Lys Thr Gly Asp Phe Ile Val
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ggc tgc ttc acc ggg ctc cag gcg gtc tcc cgg gtc acc tcc gac cgc      528
Gly Cys Phe Thr Gly Leu Gln Ala Val Ser Arg Val Thr Ser Asp Arg
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Gln Asp Leu Gly His Arg Ile Ser Val Met Trp Asn His Val Leu Pro
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agc atc gtg ccg gcg tcc atg ctg acc tgg atc gaa acc ggc gag gag      624
Ser Ile Val Pro Ala Ser Met Leu Thr Trp Ile Glu Thr Gly Glu Glu
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cgg atc ggg aag gtc gcg gcg gcg gcc gag gcc gcc gag gct gcg gag      672
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35 40 45

Val Thr Lys Gly Ala Met Tyr Phe His Phe Ala Ser Lys Glu Glu Leu
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Ala Arg Gly Val Leu Ala Glu Gln Thr Leu His Val Ala Val Pro Glu
65 70 75 80

Ser Gly Ser Lys Ala Gln Glu Leu Val Asp Leu Thr Met Leu Val Ala
85 90 95

His Gly Met Leu His Asp Pro Ile Leu Arg Ala Gly Thr Arg Leu Ala
100 105 110

Leu Asp Gln Gly Ala Val Asp Phe Ser Asp Ala Asn Pro Phe Gly Glu
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Trp Gly Asp Ile Cys Ala Gln Leu Leu Ala Glu Ala Gln Glu Arg Gly
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Gln Asp Leu Gly His Arg Ile Ser Val Met Trp Asn His Val Leu Pro
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<301> Kinoshita, H., Tsuji, T., Ipposhi, H., Nihira, T. and Yamada, Y.

<302> Characterization of Binding Sequences for Butyrolactone Autoregulator Receptors in Streptomycetes

<303> Journal of Bacteriology

<304> 181

<305> 16

<306> 5075-5080

<307> 1999-08

<308> D32251

<309> 1994-07-19

<313> (1)..(26)

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<301> Kinoshita, H., Tsuji, T., Ipposhi, H., Nihira, T. and Yamada, Y.

<302> Characterization of Binding Sequences for Butyrolactone Autoregulator Receptors in Streptomycetes

<303> Journal of Bacteriology

<304> 181

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<307> 1999-08

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<212> DNA

<213> Artificial sequence

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<223> Designed sequence of the CaMV 35S promoter modified to contain the operator BARE-3 element just upstream of its TATA-box

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<223> Designed sequence of the CaMV 35S promoter modified to contain the operator BARE-3 elements just downstream and upstream of its TATA-box

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<223> Designed sequence of the CaMV 35S promoter modified to contain three of the operator BARE-3 elements just downstream and upstream of its TATA-box

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<223> Designed sequence of a backward primer containing the restriction enzyme BamH I recognition sequence for PCR amplification of the barA gene coding region to be cloned by cut with the enzyme

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<212> DNA

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<212> DNA

<213> Artificial sequence

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atgtatctct tatatagtca aaagaaccg 89